

IN THE EUROPEAN PATENT OFFICE  
BEFORE THE INTERNATIONAL SEARCHING AUTHORITY

In re Application of: COR Therapeutics, Inc.

International Application No.:PCT/US99/17594

International Application Filed: August 4, 1999

Title: **Transgenic Animals Having a Modified Glycoprotein V Gene and Methods for Their Use**

European Patent Office  
P.B. 5818 Patentlaan 2  
NL-2280 HV Rijswijk, The Netherlands

Attn: EPO - Authorized Officer

Dear Authorized Officer:

**STATEMENT ACCOMPANYING SEQUENCE LISTING**

The undersigned hereby states upon information and belief that the Sequence Listing submitted concurrently herewith does not include matter which goes beyond the content of the application as filed and that the information recorded on the diskette submitted concurrently herewith is identical to the written Sequence Listing submitted herewith.

Respectfully submitted,

By: Rosanne Kosson  
Printed Name: Rosanne Kosson

Dated: December 15, 1999

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SEQUENCE LISTING

<110> COR Therapeutics, Inc.  
Ramakrishnan, Vanitha  
Phillips, David

<120> Transgenic Animals Having a Modified Glycoprotein V  
Gene and Methods for Their Use

<130> 44481-5044-WO

<140> PCT/US99/17594  
<141> 1999-08-04

<150> US 60/109,797  
<151> 1998-08-04

<160> 14

<170> PatentIn Ver. 2.1

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<222> (3)..(24)  
<223> y at positions 3, 12, 15, 18 and 19 = c or t; v at  
position 6 = a or c or g; m at position 9 = a or  
c; s at position 24 = c or g.  
  
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26

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<221> variation  
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b at position 4 = c or g or t; k at position 6 =  
g or t.  
  
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21

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 gagccttgg tttgcttttgcatttcagac atg cta aga agc gcc ctg ctg tcc 1434  
 Met Leu Arg Ser Ala Leu Leu Ser  
 1 5

gcg gtg ctc gca ctc ttg cgt gcc caa cct ttt ccc tgc ccc aaa acc 1482  
 Ala Val Leu Ala Leu Leu Arg Ala Gln Pro Phe Pro Cys Pro Lys Thr  
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 Cys Lys Cys Val Val Arg Asp Ala Ala Gln Cys Ser Gly Gly Ser Val  
 25 30 35 40

gct cac atc gct gag cta ggt ctg cct acg aac ctc aca cac atc ctg 1578  
 Ala His Ile Ala Glu Leu Gly Leu Pro Thr Asn Leu Thr His Ile Leu  
 45 50 55

ctc ttc cga atg gac cag ggc ata ttg cgg aac cac agc ttc agc ggc 1626  
 Leu Phe Arg Met Asp Gln Gly Ile Leu Arg Asn His Ser Phe Ser Gly  
 60 65 70

atg aca gtc ctt cag cgc ctg atg ctc tca gat agc cac att tcc gcc 1674  
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 75 80 85

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 Ile Asp Pro Gly Thr Phe Asn Asp Leu Val Lys Leu Lys Thr Leu Arg  
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 105 110 115 120

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 140 145 150

ttg aac cag aat cag ctc tct ttt ctt cct gct aac ctt ttc tcg agc 1914  
 Leu Asn Gln Asn Gln Leu Ser Phe Leu Pro Ala Asn Leu Phe Ser Ser  
 155 160 165

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Leu Pro Lys Gly Leu Leu Gly Ala Gln Val Lys Leu Glu Lys Leu Leu			
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ctc tat tca aac cag ctc acg tct gtg gat tcg ggg ctg ctg agc aac			2058
Leu Tyr Ser Asn Gln Leu Thr Ser Val Asp Ser Gly Leu Leu Ser Asn			
205	210	215	
ctg ggc gcc ctg act gag ctg cgg ctg gag cgg aat cac ctc cgc tcc			2106
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gta gcc ccg ggt gcc ttc gac cgc ctc gga aac ctg agc tcc ttg act			2154
Val Ala Pro Gly Ala Phe Asp Arg Leu Gly Asn Leu Ser Ser Leu Thr			
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cta tcc gga aac ctc ctg gag tct ctg ccc ggc ctc ttc ctt cac			2202
Leu Ser Gly Asn Leu Leu Glu Ser Leu Pro Pro Ala Leu Phe Leu His			
250	255	260	
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Val Ser Ser Val Ser Arg Leu Thr Leu Phe Glu Asn Pro Leu Glu Glu			
265	270	275	280
ctc ccg gac gtg ttg ttc ggg gag atg gcc ggc ctg cgg gag ctg tgg			2298
Leu Pro Asp Val Leu Phe Gly Glu Met Ala Gly Leu Arg Glu Leu Trp			
285	290	295	
ctg aac ggc acc cac ctg agc acg ccc gcc gct gcc ttc cgc aac			2346
Leu Asn Gly Thr His Leu Ser Thr Leu Pro Ala Ala Phe Arg Asn			
300	305	310	
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Leu Ser Gly Leu Gln Thr Leu Gly Leu Thr Arg Asn Pro Arg Leu Ser			
315	320	325	
gcg ctc ccg cgc ggc gtg ttc cag ggc cta cgg gag ctg cgc gtg ctc			2442
Ala Leu Pro Arg Gly Val Phe Gln Gly Leu Arg Glu Leu Arg Val Leu			
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gcg ctg cac acc aac gcc ctg gcg gag ctg cgg gac gac ggc ctg cgc			2490
Ala Leu His Thr Asn Ala Leu Ala Glu Leu Arg Asp Asp Ala Leu Arg			
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ggc ctc ggg cac ctg cgc cag gtg tcg ctg cgc cac aac cgg ctg cgg			2538
Gly Leu Gly His Leu Arg Gln Val Ser Leu Arg His Asn Arg Leu Arg			
365	370	375	
gcc ctg ccc cgc acg ctc ttc cgc aac ctc agc agc ctc gag agc gtg			2586
Ala Leu Pro Arg Thr Leu Phe Arg Asn Leu Ser Ser Leu Glu Ser Val			
380	385	390	
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Gln Leu Glu His Asn Gln Leu Glu Thr Leu Pro Gly Asp Val Phe Ala			

395

400

405

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 410 415 420

tgc gac tgt ggc ctg tgg ccc ttc ctc cag tgg ctg cgg cat cac ccg 2730  
 Cys Asp Cys Gly Leu Trp Pro Phe Leu Gln Trp Leu Arg His His Pro  
 425 430 435 440

gac atc ctg ggc cga gac gag ccc ccg cag tgc cgt ggc ccg gag cca 2778  
 Asp Ile Leu Gly Arg Asp Glu Pro Pro Gln Cys Arg Gly Pro Glu Pro  
 445 450 455

cgc gcc agc ctg tcg ttc tgg gag ctg ctg cag ggt gac ccg tgg tgc 2826  
 Arg Ala Ser Leu Ser Phe Trp Glu Leu Leu Gln Gly Asp Pro Trp Cys  
 460 465 470

ccg gat cct cgc agc ctg cct ctc gac cct cca acc gaa aat gct ctg 2874  
 Pro Asp Pro Arg Ser Leu Pro Leu Asp Pro Pro Thr Glu Asn Ala Leu  
 475 480 485

gaa gcc ccg gtt ccg tcc tgg ctg cct aac agc tgg cag tcc cag acg 2922  
 Glu Ala Pro Val Pro Ser Trp Leu Pro Asn Ser Trp Gln Ser Gln Thr  
 490 495 500

tgg gcc cag ctg gtg gcc agg ggt gaa agt ccc aat aac agg ctc tac 2970  
 Trp Ala Gln Leu Val Ala Arg Gly Glu Ser Pro Asn Asn Arg Leu Tyr  
 505 510 515 520

tgg ggt ctt tat att ctg ctt cta gta gcc cag gcc atc ata gcc gcg 3018  
 Trp Gly Leu Tyr Ile Leu Leu Val Ala Gln Ala Ile Ile Ala Ala  
 525 530 535

ttc atc gtg ttt gcc atg att aaa atc ggc cag ctg ttt cga aca tta 3066  
 Phe Ile Val Phe Ala Met Ile Lys Ile Gly Gln Leu Phe Arg Thr Leu  
 540 545 550

atc aga gag aag ctc ttg tta gag gca atg gga aaa tcg tgt 3108  
 Ile Arg Glu Lys Leu Leu Glu Ala Met Gly Lys Ser Cys  
 555 560 565

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35 40 45

Pro Thr Asn Leu Thr His Ile Leu Leu Phe Arg Met Asp Gln Gly Ile  
50 55 60

Leu Arg Asn His Ser Phe Ser Gly Met Thr Val Leu Gln Arg Leu Met  
65 70 75 80

Leu Ser Asp Ser His Ile Ser Ala Ile Asp Pro Gly Thr Phe Asn Asp  
85 90 95

Leu Val Lys Leu Lys Thr Leu Arg Leu Thr Arg Asn Lys Ile Ser Arg  
100 105 110

Leu Pro Arg Ala Ile Leu Asp Lys Met Val Leu Leu Glu Gln Leu Phe  
115 120 125

Leu Asp His Asn Ala Leu Arg Asp Leu Asp Gln Asn Leu Phe Gln Gln  
130 135 140

Leu Arg Asn Leu Gln Glu Leu Gly Leu Asn Gln Asn Gln Leu Ser Phe  
145 150 155 160

Leu Pro Ala Asn Leu Phe Ser Ser Leu Arg Glu Leu Lys Leu Leu Asp  
165 170 175

Leu Ser Arg Asn Asn Leu Thr His Leu Pro Lys Gly Leu Leu Gly Ala  
180 185 190

Gln Val Lys Leu Glu Lys Leu Leu Tyr Ser Asn Gln Leu Thr Ser  
195 200 205

Val Asp Ser Gly Leu Leu Ser Asn Leu Gly Ala Leu Thr Glu Leu Arg  
210 215 220

Leu Glu Arg Asn His Leu Arg Ser Val Ala Pro Gly Ala Phe Asp Arg  
225 230 235 240

Leu Gly Asn Leu Ser Ser Leu Thr Leu Ser Gly Asn Leu Leu Glu Ser  
245 250 255

Leu Pro Pro Ala Leu Phe Leu His Val Ser Ser Val Ser Arg Leu Thr  
 260 265 270  
 Leu Phe Glu Asn Pro Leu Glu Glu Leu Pro Asp Val Leu Phe Gly Glu  
 275 280 285  
 Met Ala Gly Leu Arg Glu Leu Trp Leu Asn Gly Thr His Leu Ser Thr  
 290 295 300  
 Leu Pro Ala Ala Ala Phe Arg Asn Leu Ser Gly Leu Gln Thr Leu Gly  
 305 310 315 320  
 Leu Thr Arg Asn Pro Arg Leu Ser Ala Leu Pro Arg Gly Val Phe Gln  
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 Gly Leu Arg Glu Leu Arg Val Leu Ala Leu His Thr Asn Ala Leu Ala  
 340 345 350  
 Glu Leu Arg Asp Asp Ala Leu Arg Gly Leu Gly His Leu Arg Gln Val  
 355 360 365  
 Ser Leu Arg His Asn Arg Leu Arg Ala Leu Pro Arg Thr Leu Phe Arg  
 370 375 380  
 Asn Leu Ser Ser Leu Glu Ser Val Gln Leu Glu His Asn Gln Leu Glu  
 385 390 395 400  
 Thr Leu Pro Gly Asp Val Phe Ala Ala Leu Pro Gln Leu Thr Gln Val  
 405 410 415  
 Leu Leu Gly His Asn Pro Trp Leu Cys Asp Cys Gly Leu Trp Pro Phe  
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 435 440 445  
 Pro Gln Cys Arg Gly Pro Glu Pro Arg Ala Ser Leu Ser Phe Trp Glu  
 450 455 460  
 Leu Leu Gln Gly Asp Pro Trp Cys Pro Asp Pro Arg Ser Leu Pro Leu  
 465 470 475 480  
 Asp Pro Pro Thr Glu Asn Ala Leu Glu Ala Pro Val Pro Ser Trp Leu  
 485 490 495  
 Pro Asn Ser Trp Gln Ser Gln Thr Trp Ala Gln Leu Val Ala Arg Gly  
 500 505 510  
 Glu Ser Pro Asn Asn Arg Leu Tyr Trp Gly Leu Tyr Ile Leu Leu Leu  
 515 520 525  
 Val Ala Gln Ala Ile Ile Ala Ala Phe Ile Val Phe Ala Met Ile Lys  
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 Ile Gly Gln Leu Phe Arg Thr Leu Ile Arg Glu Lys Leu Leu Leu Glu  
 545 550 555 560





gga atg ggc cgc ggc gtc ctg cag agc cag agc ttc agc ggc atg acc Gly Met Gly Arg Gly Val Leu Gln Ser Gln Ser Phe Ser Gly Met Thr 60	65	70	2643
gtc ctg cag cgc ctc atg atc tcc gac agc cac att tcc gcc gtt gcc Val Leu Gln Arg Leu Met Ile Ser Asp Ser His Ile Ser Ala Val Ala 75	80	85	2691
ccc ggc acc ttc agt gac ctg ata aaa ctg aaa acc ctg agg ctg tcg Pro Gly Thr Phe Ser Asp Leu Ile Lys Leu Lys Thr Leu Arg Leu Ser 95	100	105	2739
cgc aac aaa atc acg cat ctt cca ggt gcg ctg ctg gat aag atg gtg Arg Asn Lys Ile Thr His Leu Pro Gly Ala Leu Leu Asp Lys Met Val 110	115	120	2787
ctc ctg gag cag ttg ttt ttg gac cac aat gcg cta agg ggc att gac Leu Leu Glu Gln Leu Phe Leu Asp His Asn Ala Leu Arg Gly Ile Asp 125	130	135	2835
caa aac atg ttt cag aaa ctg gtt aac ctg cag gag ctc gct ctg aac Gln Asn Met Phe Gln Lys Leu Val Asn Leu Gln Glu Leu Ala Leu Asn 140	145	150	2883
cag aat cag ctc gat ttc ctt cct gcc agt ctc ttc acg aat ctg gag Gln Asn Gln Leu Asp Phe Leu Pro Ala Ser Leu Phe Thr Asn Leu Glu 155	160	165	2931
aac ctg aag ttg ttg gat tta tcg gga aac aac ctg acc cac ctg ccc Asn Leu Lys Leu Leu Asp Leu Ser Gly Asn Asn Leu Thr His Leu Pro 175	180	185	2979
aag ggg ttg ctt gga gca cag gct aag ctc gag aga ctt ctg ctc cac Lys Gly Leu Leu Gly Ala Gln Ala Lys Leu Glu Arg Leu Leu His 190	195	200	3027
tcg aac cgc ctt gtg tct ctg gat tcg ggg ctg ttg aac agc ctg ggc Ser Asn Arg Leu Val Ser Leu Asp Ser Gly Leu Leu Asn Ser Leu Gly 205	210	215	3075
gcc ctg acg gag ctg cag ttc cac cga aat cac atc cgt tcc atc gca Ala Leu Thr Glu Leu Gln Phe His Arg Asn His Ile Arg Ser Ile Ala 220	225	230	3123
ccc ggg gcc ttc gac cgg ctc cca aac ctc agt tct ttg acg ctt tcg Pro Gly Ala Phe Asp Arg Leu Pro Asn Leu Ser Ser Leu Thr Leu Ser 235	240	245	3171
aga aac cac ctt gcg ttt ctc ccc tct gcg ctc ttt ctt cat tcg cac Arg Asn His Leu Ala Phe Leu Pro Ser Ala Leu Phe Leu His Ser His 255	260	265	3219
aat ctg act ctg ttg act ctg ttc gag aac ccg ctg gca gag ctc ccg Asn Leu Thr Leu Leu Thr Leu Phe Glu Asn Pro Leu Ala Glu Leu Pro 270	275	280	3267

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Gly Val Leu Phe Gly Glu Met Gly Gly Leu Gln Glu Leu Trp Leu Asn		
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Arg Thr Gln Leu Arg Thr Leu Pro Ala Ala Ala Phe Arg Asn Leu Ser		
300	305	310
cgc ctg cgg tac tta ggg gtg act ctg agc cgg cgg ctg agc gcg ctt		3411
Arg Leu Arg Tyr Leu Gly Val Thr Leu Ser Pro Arg Leu Ser Ala Leu		
315	320	325
ccg cag ggc gcc ttc cag ggc ctt ggc gag ctc cag gtg ctc gcc ctg		3459
Pro Gln Gly Ala Phe Gln Gly Leu Gly Glu Leu Gln Val Leu Ala Leu		
335	340	345
cac tcc aac ggc ctg acc gcc ctc ccc gac ggc ttg ctg cgc ggc ctc		3507
His Ser Asn Gly Leu Thr Ala Leu Pro Asp Gly Leu Leu Arg Gly Leu		
350	355	360
ggc aag ctg cgc cag gtg tcc ctg cgc cgc aac agg ctg cgc gcc ctg		3555
Gly Lys Leu Arg Gln Val Ser Leu Arg Arg Asn Arg Leu Arg Ala Leu		
365	370	375
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395	400	410
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Pro Arg Leu Thr Glu Val Leu Leu Gly His Asn Ser Trp Arg Cys Asp		
415	420	425
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Cys Gly Leu Gly Pro Phe Leu Gly Trp Leu Arg Gln His Leu Gly Leu		
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Val Gly Gly Glu Glu Pro Pro Arg Cys Ala Gly Pro Gly Ala His Ala		
445	450	455
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Gly Leu Pro Leu Trp Ala Leu Pro Gly Gly Asp Ala Glu Cys Pro Gly		
460	465	470
ccc cgg ggc ccc cct ccc cgc ccc gct ggc gac agc tcc tcg gaa gcc		3891
Pro Arg Gly Pro Pro Pro Arg Pro Ala Ala Asp Ser Ser Glu Ala		
475	480	490
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Pro Val His Pro Ala Leu Ala Pro Asn Ser Ser Glu Pro Trp Val Trp		
495	500	505

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Ala Gln Pro Val Thr Thr Gly Lys Gly Gln Asp His Ser Pro Phe Trp	
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Leu Gln Ser Gln Ser Phe Ser Gly Met Thr Val Leu Gln Arg Leu Met  
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Ile Ser Asp Ser His Ile Ser Ala Val Ala Pro Gly Thr Phe Ser Asp  
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Leu Ile Lys Leu Lys Thr Leu Arg Leu Ser Arg Asn Lys Ile Thr His  
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Gln Ala Lys Leu Glu Arg Leu Leu Leu His Ser Asn Arg Leu Val Ser  
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Thr Leu Pro Gly Asp Val Phe Gly Ala Leu Pro Arg Leu Thr Glu Val  
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Pro Arg Cys Ala Gly Pro Gly Ala His Ala Gly Leu Pro Leu Trp Ala  
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Gly Lys Gly Gln Asp His Ser Pro Phe Trp Gly Phe Tyr Phe Leu Leu  
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